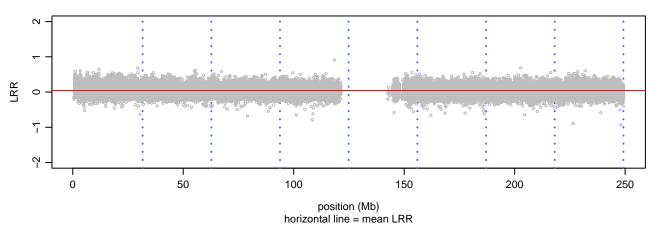
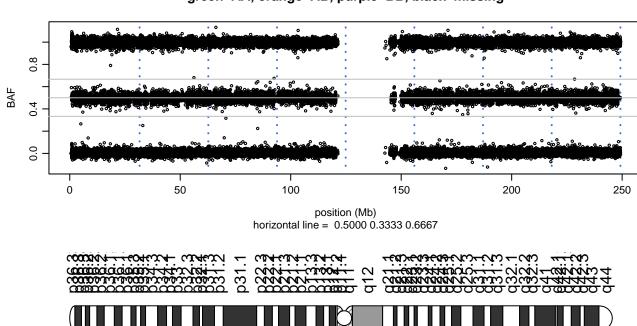
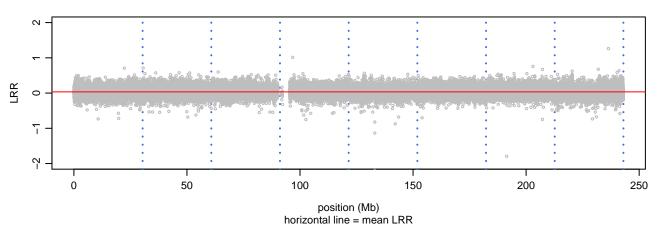
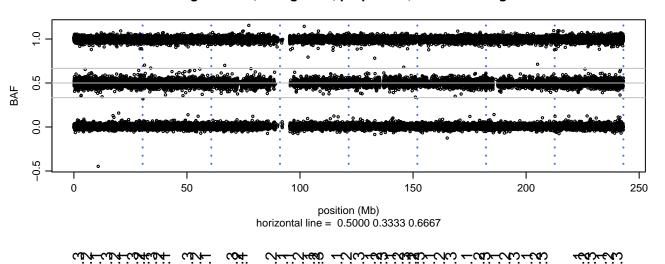
Scan 280 - Chromosome 1 - KCNA1_R417X_C01



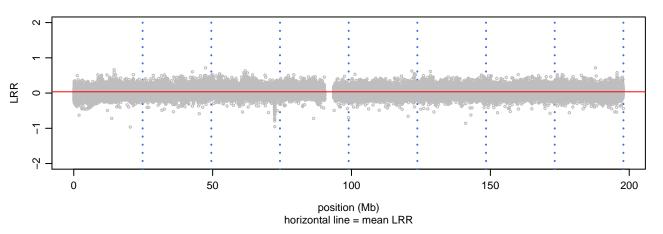


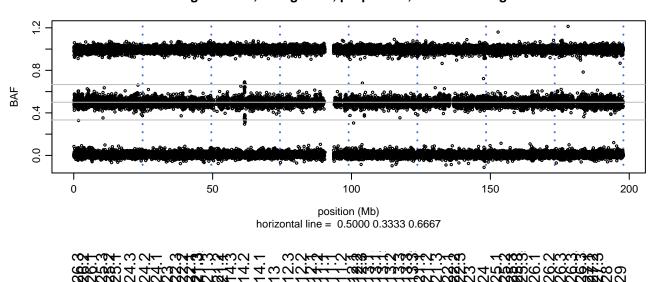
Scan 280 - Chromosome 2 - KCNA1_R417X_C01



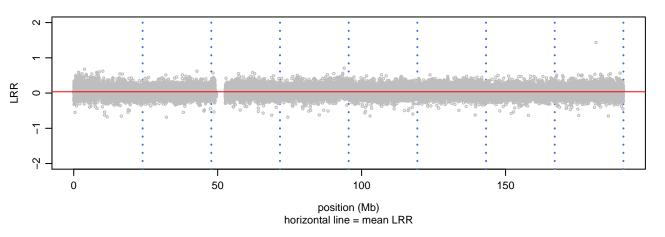


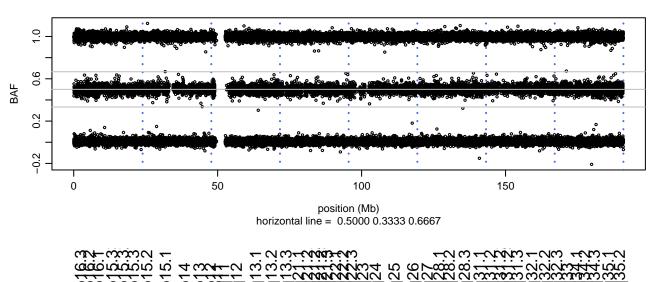
Scan 280 - Chromosome 3 - KCNA1_R417X_C01



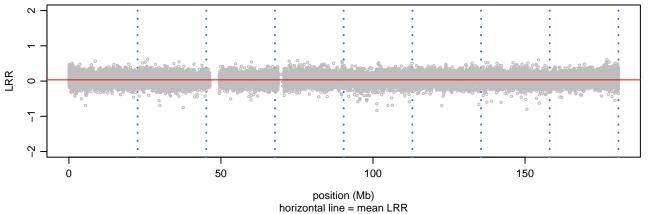


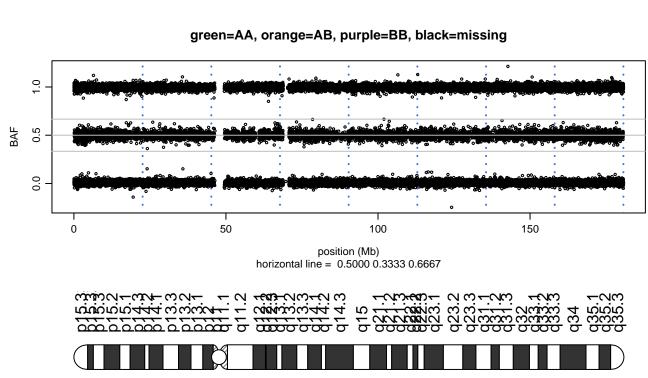
Scan 280 - Chromosome 4 - KCNA1_R417X_C01



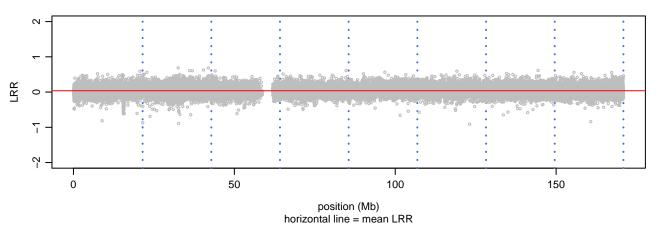


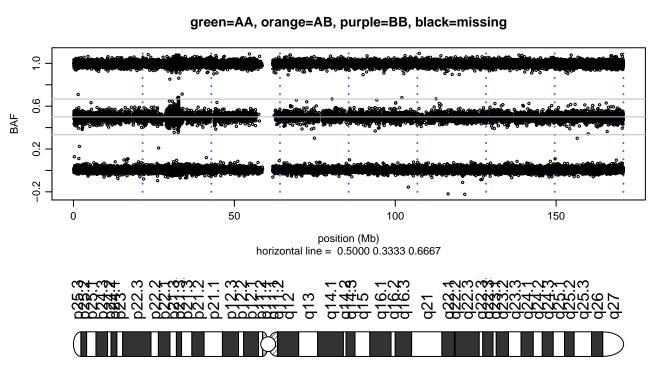
Scan 280 - Chromosome 5 - KCNA1_R417X_C01



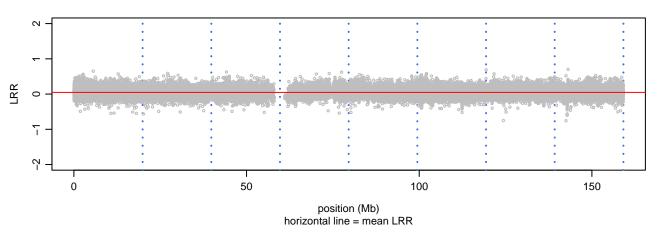


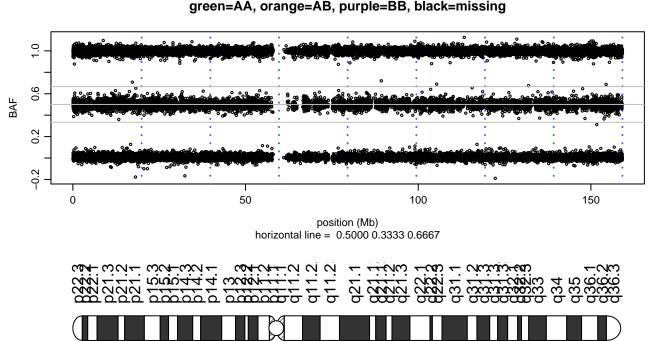
Scan 280 - Chromosome 6 - KCNA1_R417X_C01



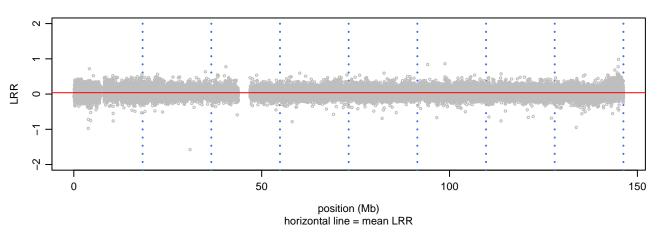


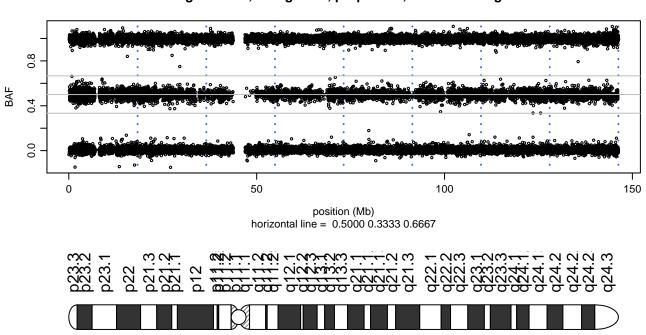
Scan 280 - Chromosome 7 - KCNA1_R417X_C01



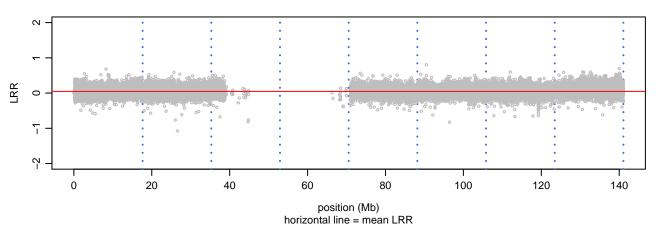


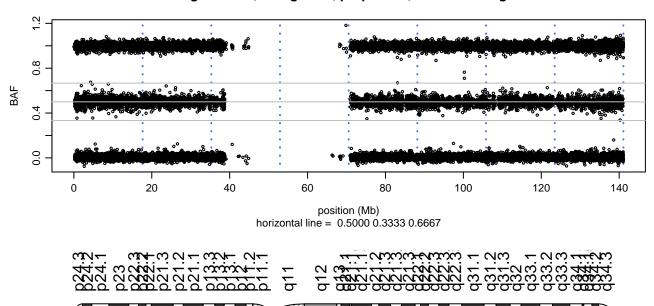
Scan 280 - Chromosome 8 - KCNA1_R417X_C01



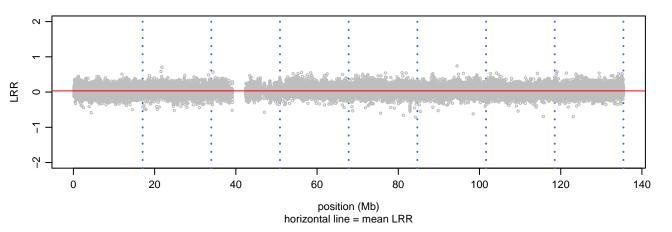


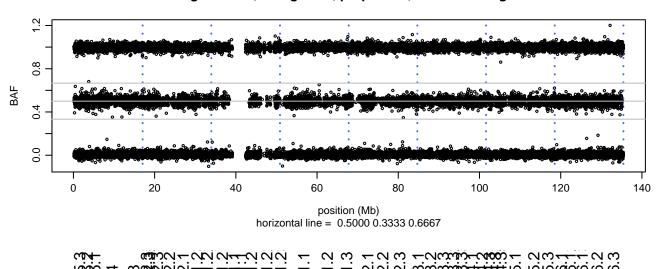
Scan 280 - Chromosome 9 - KCNA1_R417X_C01



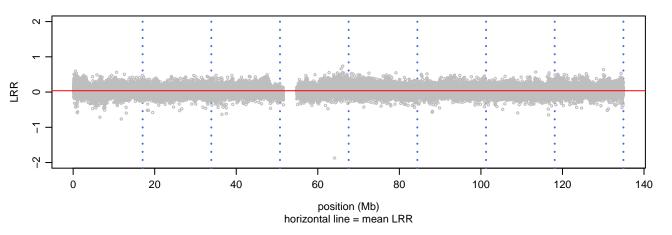


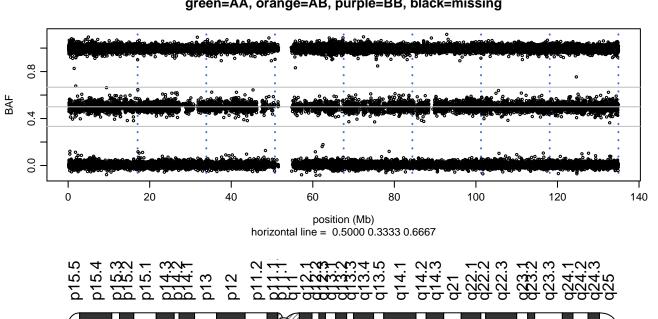
Scan 280 - Chromosome 10 - KCNA1_R417X_C01



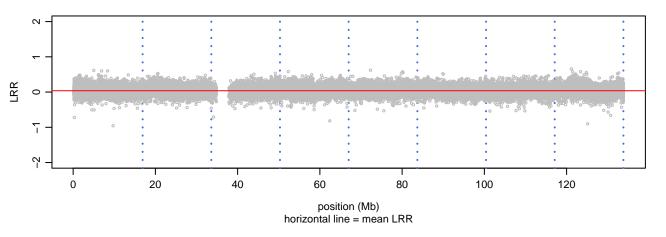


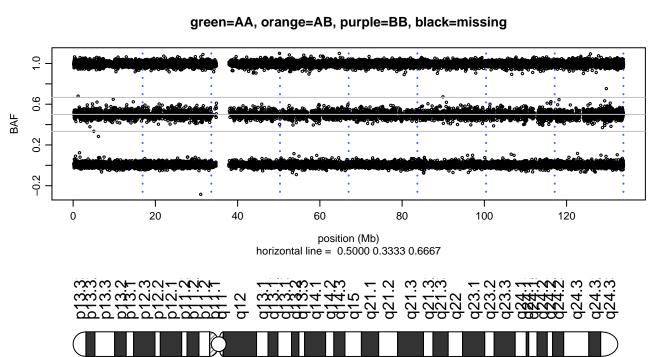
Scan 280 - Chromosome 11 - KCNA1_R417X_C01



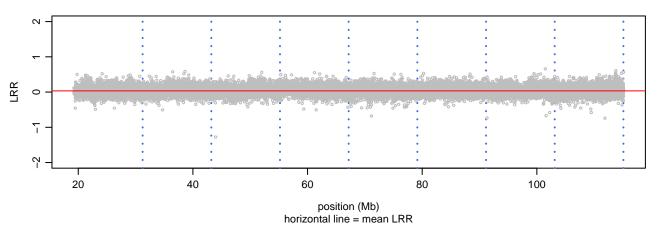


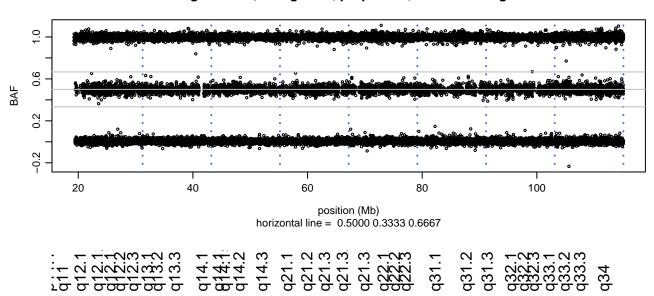
Scan 280 - Chromosome 12 - KCNA1_R417X_C01



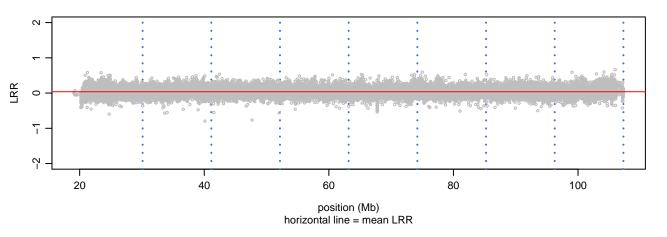


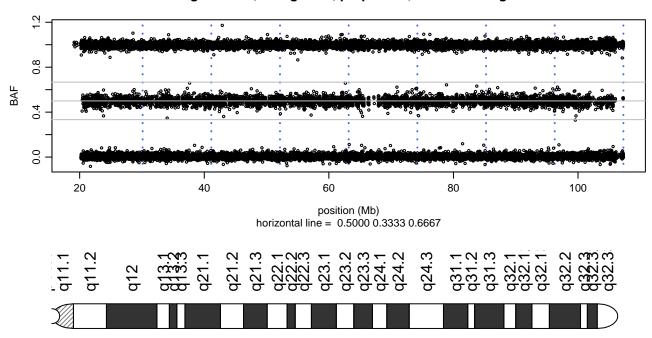
Scan 280 - Chromosome 13 - KCNA1_R417X_C01



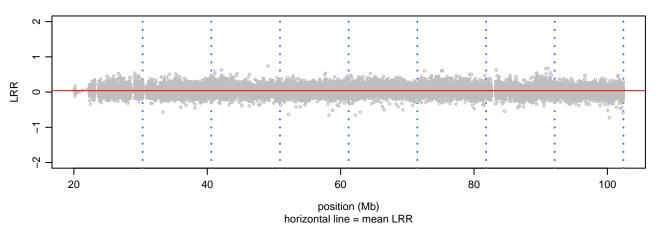


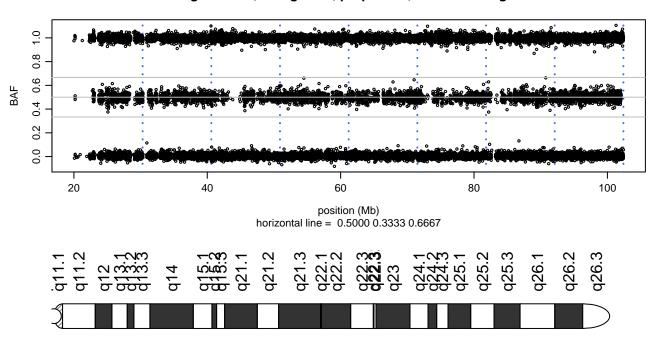
Scan 280 - Chromosome 14 - KCNA1_R417X_C01



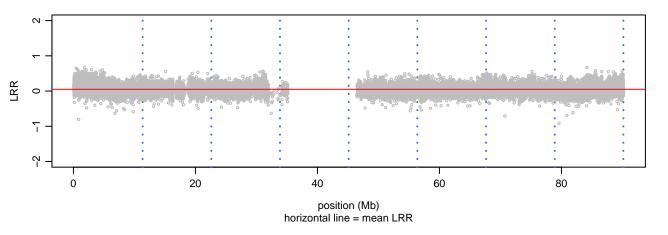


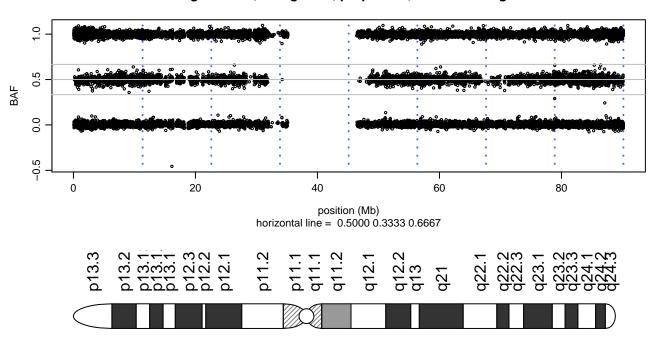
Scan 280 - Chromosome 15 - KCNA1_R417X_C01



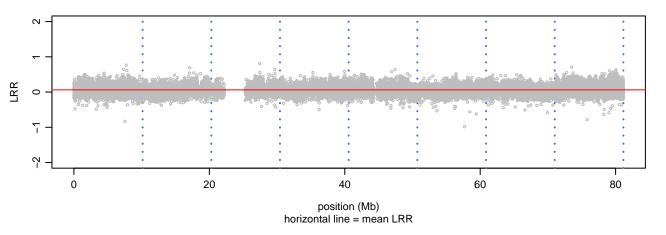


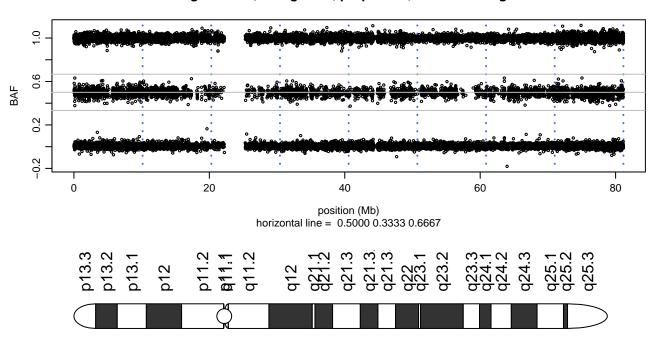
Scan 280 - Chromosome 16 - KCNA1_R417X_C01



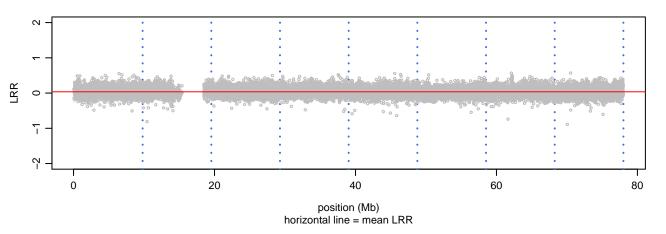


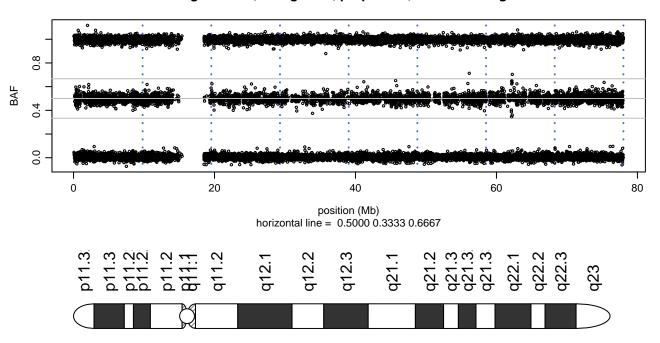
Scan 280 - Chromosome 17 - KCNA1_R417X_C01



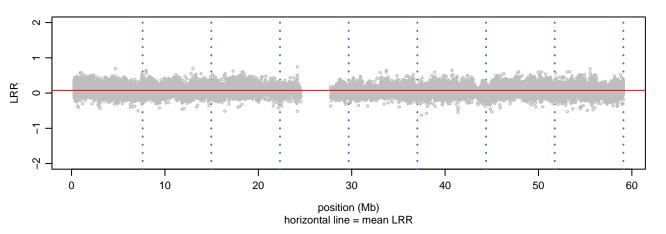


Scan 280 - Chromosome 18 - KCNA1_R417X_C01

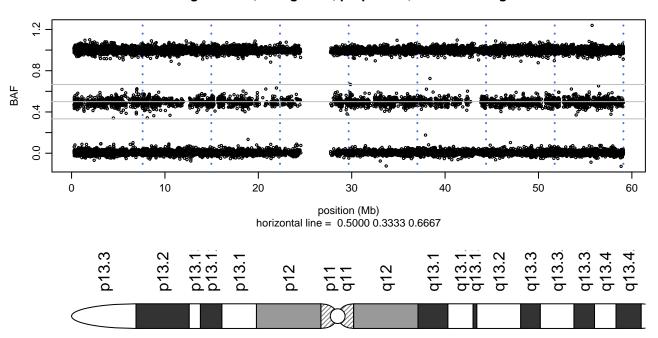




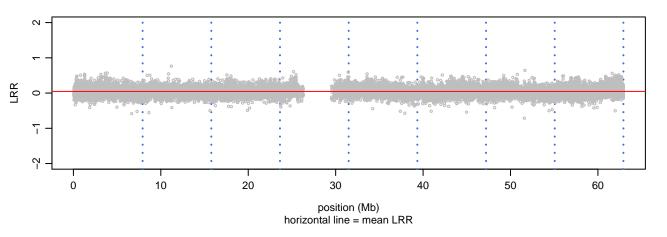
Scan 280 - Chromosome 19 - KCNA1_R417X_C01

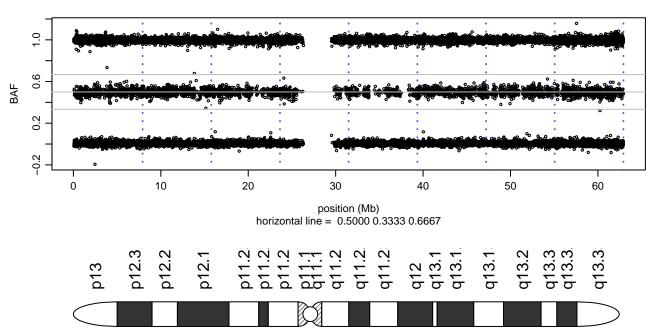


green=AA, orange=AB, purple=BB, black=missing

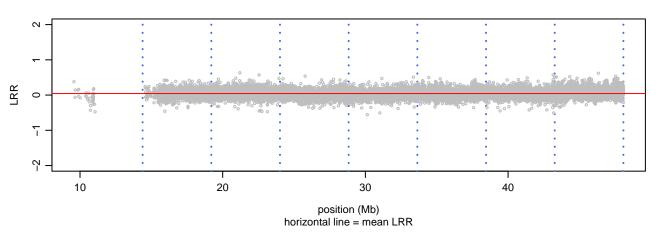


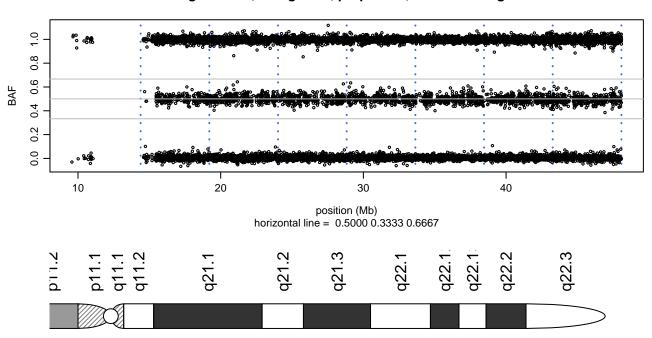
Scan 280 - Chromosome 20 - KCNA1_R417X_C01



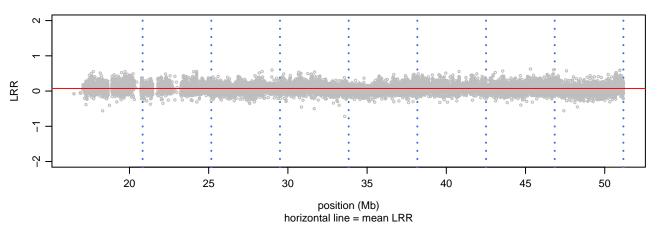


Scan 280 - Chromosome 21 - KCNA1_R417X_C01

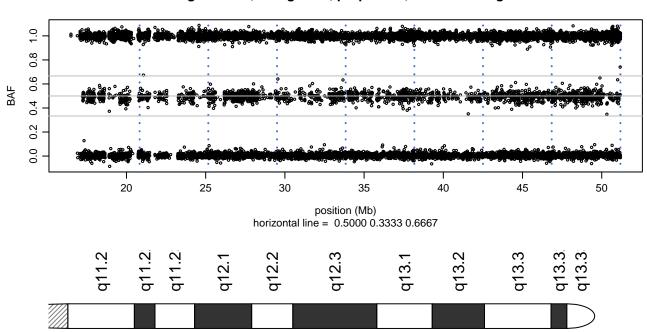




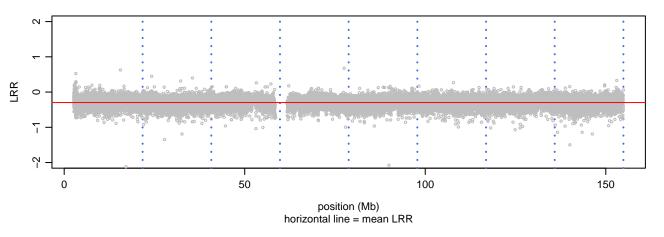
Scan 280 - Chromosome 22 - KCNA1_R417X_C01

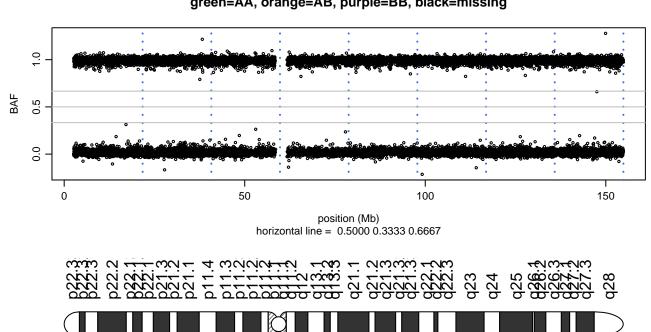


green=AA, orange=AB, purple=BB, black=missing

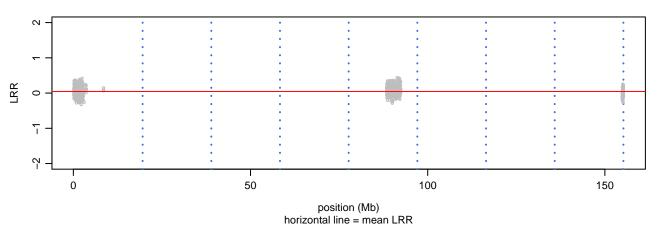


Scan 280 - Chromosome X - KCNA1_R417X_C01

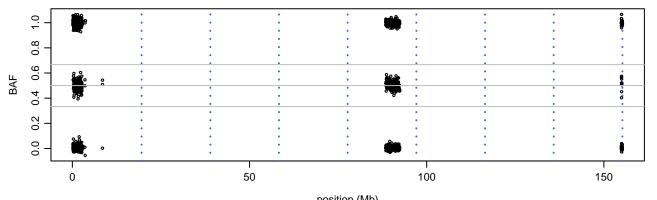




Scan 280 - Chromosome Y - KCNA1_R417X_C01



green=AA, orange=AB, purple=BB, black=missing



position (Mb) horizontal line = 0.5000 0.3333 0.6667

